

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/516,444A

Input Set : A:\5995USP2.txt

Output Set: N:\CRF4\08082002\I516444A.raw

```
<110> APPLICANT: Abbott Laboratories
              Billing-Medel, Patricia A.
      5
              Cohen, Maurice
      6
     7
              Colpitts, Tracey L.
      8
              Friedman, Paula N.
      9
              Gordon, Julian
              Granados, Edward N.
     10
              Hodges, Steven C.
     11
     12
              Klass, Michael R.
     13
              Kratochvil, Jon D.
              Roberts-Rapp, Lisa
     14
     15
              Russell, John C.
              Stroupe, Stephen D.
     18 <120> TITLE OF INVENTION: Reagents And Method Useful For Detecting
              Diseases Of The Breast
     19
     22 <130> FILE REFERENCE: 5995.US.P2
     24 <140> CURRENT APPLICATION NUMBER: 09/516,444A
     25 <141> CURRENT FILING DATE: 2000-02-29
     27 <150> PRIOR APPLICATION NUMBER: US 08/962,094
     28 <151> PRIOR FILING DATE: 1997-10-31
     30 <150> PRIOR APPLICATION NUMBER: US 08/742,067
     31 <151> PRIOR FILING DATE: 1996-10-31
     33 <160> NUMBER OF SEQ ID NOS: 39
     35 <170> SOFTWARE: FastSEQ for Windows Version 4.0
     37 <210> SEQ ID NO: 1
     38 <211> LENGTH: 201
     39 <212> TYPE: DNA
     40 <213> ORGANISM: Artificial Sequence
     42 <220> FEATURE:
     43 <223> OTHER INFORMATION: EST Clone 1662885
W--> 45 <221> NAME/KEY: misc_feature
     46 <222> LOCATION: (26)...(26)
     47 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other at
     48
              position 26
W--> 50 <221> misc_feature
     51 <222> LOCATION: (98)...(98)
     52 < 223 > OTHER INFORMATION: n = a or g or c or t/u, unknown or other at
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W--> 55 <221> misc_feature
     56 <222> LOCATION: (133)...(133)
     57 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other at
     58
              position 133
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W--> 60 <221> misc_feature

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DATE: 08/08/2002 \stackrel{\frown}{\mathbf{P}}
                     RAW SEQUENCE LISTING
                                                               TIME: 12:03:45
                     PATENT APPLICATION: US/09/516,444A
                     Input Set : A:\5995USP2.txt
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     61 <222> LOCATION: (145)...(145)
     62 < 223 > OTHER INFORMATION: n = a or g or c or t/u, unknown or other at
     63
              position 145
W--> 65 <221> misc_feature
     66 <222> LOCATION: (183)...(183)
     67 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other at
              position 183
W--> 70 < 400 > 1
W--> 71 ctcttaggct ttgaagcatt tttgtntgtg ctccctgatc ttcatgtcac caccatgaag 60
W--> 72 ttcttagcag tcctggtact cttgggagtt tccatctntc tggtctctgc ccagaatccg 120
W--> 73 acaacagctg ctncagctga cacgnatcca gctactggtc ctgctgatga tgaagcccct 180
W--> 74 gangctgaaa ccactgctgc t
     76 <210> SEQ ID NO: 2
     77 <211> LENGTH: 308
     78 <212> TYPE: DNA
     79 <213> ORGANISM: Homo sapiens
     81 <400> SEQUENCE: 2
     82 taggetttga ageatttttg tetgtgetee etgatettea ggteaceace atgaagttet 60
     83 tagcagteet ggtactettg ggagttteea tetttetggt etetgeecag aateegacaa 120
     84 cagetgetee agetgacaeg tatecageta etggteetge tgatgatgaa geceetgatg 180
     85 ctgaaaccac tgctgctgca accactgcga ccactgctgc tcctaccact gcaaccaccg 240
     86 ctgcttctac cactgctcgt aaagacattc cagttttacc caaatgggtt ggggatcttc 300
     87 cgaatggt
     89 <210> SEO ID NO: 3
     90 <211> LENGTH: 292
     91 <212> TYPE: DNA
     92 <213> ORGANISM: Artificial Sequence
     94 <220> FEATURE:
     95 <223> OTHER INFORMATION: EST Clone 901429
W--> 97 <221> NAME/KEY: misc_feature
     98 <222> LOCATION: (236)...(236)
     99 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other at
               position 236
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W--> 102 <221> misc_feature
     103 <222> LOCATION: (259)...(259)
     104 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other at
               position 259
W--> 107 <400> 3
     108 gcatttttgt ctgtgctccc tgatcttcat gtcaccacca tgaagttctt agcagtcctg 60
     109 gtactettgg gagtttecat etttetggte tetgeceaga ateegacaac agetgeteea 120
     110 gctgacacgt atccagctac tggtcctgct gatgatgaag cccctgatgc tgaaaccact 180
W--> 111 gctgctgcaa ccactgcgac cactgctgct cctaccactg caaccaccgc tgcttntacc 240
W--> 112 actgctcgta aagacattnc agttttaccc aaatgggttg gggatctccc ga
     114 <210> SEQ ID NO: 4
     115 <211> LENGTH: 197
     116 <212> TYPE: DNA
     117 <213> ORGANISM: Homo sapiens
     119 <400> SEQUENCE: 4
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120 gttttaccca aatgggttgg ggatctcccg aatggtagag tgtgtccctg agatggaatc 60

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121 agettgagte ttetgeaatt ggteacaact atteatgett eetgtgattt eateeaacta 120 122 cttaccttgc ctacgatatc ccctttatct ctaatcagtt tattttcttt caaataaaaa 180 123 ataactatga gcaacat 125 <210> SEQ ID NO: 5 126 <211> LENGTH: 472 127 <212> TYPE: DNA 128 <213> ORGANISM: Homo sapiens 130 <400> SEQUENCE: 5 131 ctcttagget ttgaageatt tttgtctgtg ctccctgate ttcatgtcac caccatgaag 60 132 ttcttagcag tcctggtact cttgggagtt tccatctttc tggtctctgc ccagaatccg 120 133 acaacagctg ctccagctga cacgtatcca gctactggtc ctgctgatga tgaagcccct 180 134 gatgetgaaa ceaetgetge tgeaaceaet gegaeeaetg etgeteetae eaetgeaaee 240 135 acceptgett ctaccactge tegtaaagac attecagttt tacccaaatg ggttggggat 300 136 ctcccgaatg gtagagtgtg tccctgagat ggaatcagct tgagtcttct gcaattggtc 360 137 acaactatte atgetteetg tgattteate caactaetta cettgeetae gatateeeet 420 138 ttatctctaa tcagtttatt ttctttcaaa taaaaaataa ctatgagcaa ca 140 <210> SEQ ID NO: 6 141 <211> LENGTH: 473 142 <212> TYPE: DNA 143 <213> ORGANISM: Homo sapiens 145 <400> SEQUENCE: 6 146 ctcttaggct ttgaagcatt tttgtctgtg ctccctgatc ttcatgtcac caccatgaag 60 147 ttettageag teetggtact ettgggagtt teeatettte tggtetetge ceagaateeg 120 148 acaacagctg ctccagctga cacgtatcca gctactggtc ctgctgatga tgaagcccct 180 149 gatgctgaaa ccactgctgc tgcaaccact gcgaccactg ctgctcctac cactgcaacc 240 150 acceptett ctaccactec testaaagac attecasttt tacccaaats ggttggggat 300 151 ctcccgaatg gtagagtgtg tccctgagat ggaatcagct tgagtcttct gcaattggtc 360 152 acaactatte atgetteetg tgattteate caactaetta cettgeetae gatateeeet 420 153 ttatctctaa tcagtttatt ttctttcaaa taaaaaataa ctatgagcaa cat 155 <210> SEQ ID NO: 7 156 <211> LENGTH: 68 157 <212> TYPE: DNA 158 <213> ORGANISM: Artificial Sequence 160 <220> FEATURE: 161 <223> OTHER INFORMATION: Restriction site 163 <400> SEQUENCE: 7 164 agctcggaat tccgagcttg gatcctctag agcggccgcc gactagtgag ctcgtcgacc 60 165 cgggaatt 167 <210> SEQ ID NO: 8 168 <211> LENGTH: 68 169 <212> TYPE: DNA 170 <213> ORGANISM: Artificial Sequence 172 <220> FEATURE: 173 <223> OTHER INFORMATION: Restriction site 175 <400> SEQUENCE: 8 176 aattaattcc cgggtcgacg agctcactag tcggcggccg ctctagagga tccaagctcg 60 68 177 gaattccg 179 <210> SEQ ID NO: 9 180 <211> LENGTH: 24

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181 <212> TYPE: DNA 182 <213> ORGANISM: Artificial Sequence 184 <220> FEATURE: 185 <223> OTHER INFORMATION: Universal Primer 187 <400> SEQUENCE: 9 24 188 agcggataac aatttcacac agga 190 <210> SEQ ID NO: 10 191 <211> LENGTH: 18 192 <212> TYPE: DNA 193 <213> ORGANISM: Artificial Sequence 195 <220> FEATURE: 196 <223> OTHER INFORMATION: Universal Primer 198 <400> SEQUENCE: 10 18 199 tgtaaaacga cggccagt 201 <210> SEQ ID NO: 11 202 <211> LENGTH: 20 203 <212> TYPE: DNA 204 <213> ORGANISM: Artificial Sequence 206 <220> FEATURE: 207 <223> OTHER INFORMATION: Primer 209 <400> SEQUENCE: 11 20 210 actgctcgta aagacattcc 212 <210> SEQ ID NO: 12 213 <211> LENGTH: 19 214 <212> TYPE: DNA 215 <213> ORGANISM: Artificial Sequence 217 <220> FEATURE: 218 <223> OTHER INFORMATION: Primer 220 <400> SEQUENCE: 12 19 221 gggacacact ctaccattc 223 <210> SEQ ID NO: 13 224 <211> LENGTH: 20 225 <212> TYPE: DNA 226 <213> ORGANISM: Artificial Sequence 228 <220> FEATURE: 229 <223> OTHER INFORMATION: Sense Primer 231 <400> SEQUENCE: 13 20 232 aagcccctga tgctgaaacc 234 <210> SEQ ID NO: 14 235 <211> LENGTH: 23 236 <212> TYPE: DNA 237 <213> ORGANISM: Artificial Sequence 239 <220> FEATURE: 240 <223> OTHER INFORMATION: Antisense Primer 242 <400> SEQUENCE: 14 23 243 tgcagaagac tcaagctgat tcc 245 <210> SEQ ID NO: 15 246 <211> LENGTH: 20

247 <212> TYPE: DNA





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Input Set : A:\5995USP2.txt

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	213> ORGANISM: Artificial Sequence	
	220> FEATURE: 223> OTHER INFORMATION: Target-Specific Forward Primer	
	400> SEQUENCE: 15	
	agcecetga tgetgaaace	20
	210> SEQ ID NO: 16	
	211> LENGTH: 23	
	212> TYPE: DNA	
	213> ORGANISM: Artificial Sequence	
	220> FEATURE:	
262	223> OTHER INFORMATION: Target-Specific Reverse Primer	
264	400> SEQUENCE: 16	
265	gcagaagac tcaagctgat tcc	23
267	210> SEQ ID NO: 17	
	211> LENGTH: 15	
	212> TYPE: DNA	
	213> ORGANISM: Artificial Sequence	
	220> FEATURE:	
	223> OTHER INFORMATION: Probe	
	400> SEQUENCE: 17	1 =
	accactgct gctcc	15
	210> SEQ ID NO: 18	
	211> LENGTH: 20	
	212> TYPE: DNA	
	213> ORGANISM: Artificial Sequence 220> FEATURE:	
	223> OTHER INFORMATION: Sense Primer	
	400> SEQUENCE: 18	
	ctgctcgta aagacattcc	20
	210> SEQ ID NO: 19	
	211> LENGTH: 19	
	212> TYPE: DNA	
292	213> ORGANISM: Artificial Sequence	
	220> FEATURE:	
295	223> OTHER INFORMATION: Antisense Primer	
297	400> SEQUENCE: 19	
298	ggacacact ctaccattc	19
300	210> SEQ ID NO: 20	
	211> LENGTH: 90	
302	212> TYPE: PRT	
	213> ORGANISM: Homo sapiens	
	400> SEQUENCE: 20	
	et Lys Phe Leu Ala Val Leu Val Leu Gly Val Ser Ile Phe Leu	
307		
	al Ser Ala Gln Asn Pro Thr Thr Ala Ala Pro Ala Asp Thr Tyr Pro	
309	20 25 30	
	la Thr Gly Pro Ala Asp Asp Glu Ala Pro Asp Ala Glu Thr Thr Ala 35 40 45	
311	35 40 45 la Ala Thr Thr Ala Thr Thr Ala Ala Pro Thr Thr Ala Thr Thr Ala	
2TZ	ta wia int lift wia int lift wia wia 210 int int wia int lift wia	

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/516,444A

DATE: 08/08/2002 TIME: 12:03:46

Input Set : A:\5995USP2.txt

Output Set: N:\CRF4\08082002\I516444A.raw

L:45 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:50 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:55 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:60 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:65 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:70 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:71 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0
L:72 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:60
L:73 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:120
L:74 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:180
L:97 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:102 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:3
L:107 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:3
L:111 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:180
L:112 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:180
L:112 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:240



Creation date: 11-14-2003

Indexing Officer: QNGUYEN5 - QUANG NGUYEN

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Dossier: 09516444

Legal Date: 10-02-2003

No.	Doccode	Number of pages
1	CTRS	6
2	NFDR	1
3	EXIN	1

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Remarks:

Order of re-scan issued on